

## Genotyping protocol

Project Ncam2

(PHENOMIN-ICS reference IR00006055 / P6055)

This report has been **prepared** by: Christelle MORGENTHALER-ROTH

This report has been **validated** by: Sylvie Jacquot, PhD  
Head of Genotyping Service

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**For any question, please contact:**

**PHENOMIN-ICS**

Email: [genotypingrequest@igbmc.fr](mailto:genotypingrequest@igbmc.fr)

Web site: <http://www.ics-mci.fr/>



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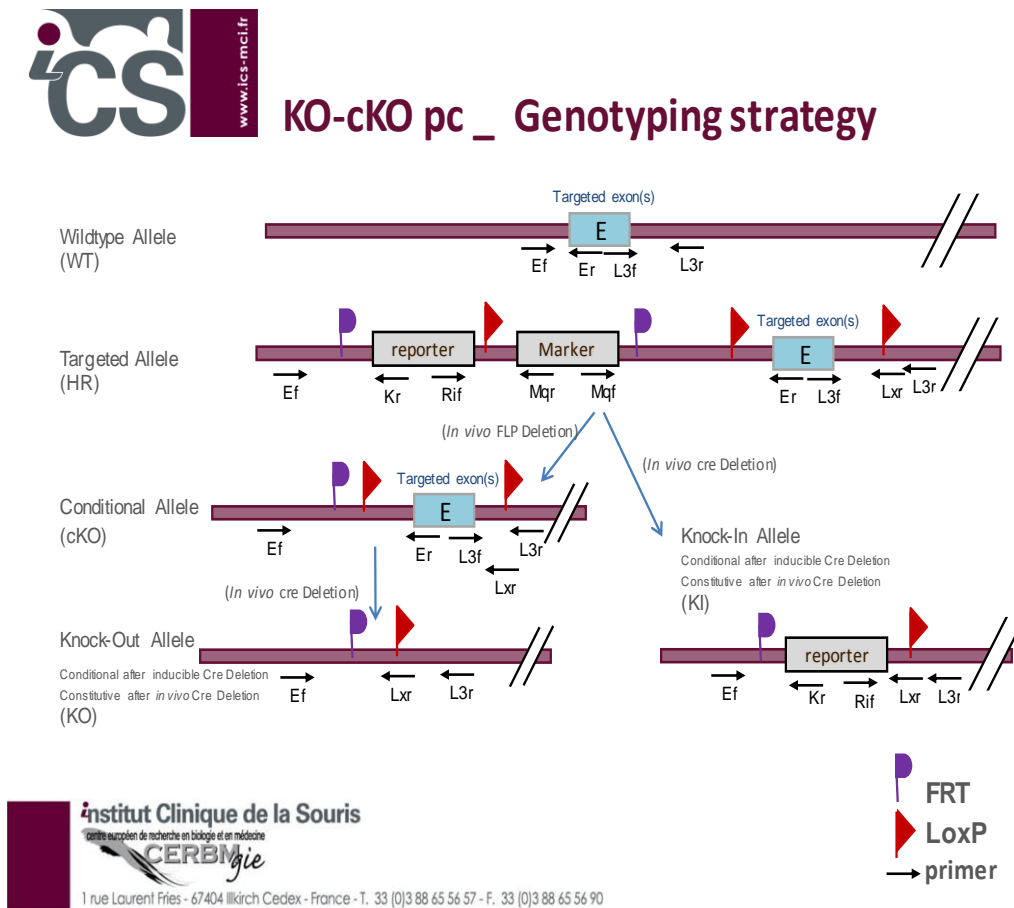


## 1. Genotyping protocol and data

This section describes the condition used at the Mouse Clinical Institute (ICS) to genotype your **Ncam2** Constitutive Knockout / Conditional Knockout (KO-cKO x Cre) project.

### 1.1. Genotyping strategy

The map below describes the position of the primers used for genotyping for each possible allele.



Sequence of primers used for genotyping:

| Position | Primers | Sequence                        |
|----------|---------|---------------------------------|
| Ef       | 9317    | GGGGAAAGAAGCTTGGAAATGATAATTACTG |
| Er       | 9320    | GTCAAGCAACCTTAAGGCAAACACATAG    |
| Kr       | 3277    | CTCCTACATAGTTGGCAGTGTGGG        |
| L3f      | 9318    | CCAAGTGTAAGGATGTCAGAGCCAACC     |
| L3r      | 9319    | CAACTGTCTTTTCATTGTCAAACGAAGAG   |
| Lxr      | 3255    | ACTGATGGCGAGCTCAGACCATAAC       |
| Rif      | 5966    | GCACATGGCTGAATATCGACGGT         |

PCR fragments expected size (bp):

| Region analyzed                                 | Primers used | Position on the primer<br>(see the map above) | Targeted allele (HR) | conditional allele (cKO) | KI allele | WildType allele |
|---|--------------|---|----------------------|--------------------------|-----------|-----------------|
| 5' part of the selection marker (with Betaine)  | 9317-3277    | Ef / Kr                                       | 274                  | ---                      | ---       | ---             |
| Presence of the distal loxP                     | 9318-9319    | L3f / L3r                                     | 385                  | 385                      | ---       | 408             |
| Distal loxP specific PCR                        | 9318-3255    | L3f / Lxr                                     | 247                  | 247                      | ---       | ---             |
| Excision of the selection marker (with Betaine) | 9317-9320    | Ef / Er                                       | 7464*                | 560                      | ---       | 410             |
| Cre total excision                              | 5966-3255    | Rif / Lxr                                     | 3255*                | ---                      | 471       | ---             |

\*: this PCR product will not be observed using our PCR genotyping conditions (see description below)

---: no Amplicon should be obtained



## 1.2. PCR protocol

This section describes the composition of the mix and cycling conditions used for genotyping.

| Reagents:                      | Volume:     |
|--------------------------------|-------------|
| - FastStart PCR Master (Roche) | 7.5µl       |
| - DNA (50ng/µl)                | 1.5µl       |
| - 5' primer (100 µM)           | 0.06µl      |
| - 3' primer (100 µM)           | 0.06µl      |
| - Sterile H <sub>2</sub> O     | up to 15 µl |

### Cycling conditions:

| Temp | Time | #Cycles |
|------|------|---------|
| 95°C | 4min | 1       |
| 94°C | 30s  | 34      |
| 62°C | 30s  |         |
| 72°C | 1min |         |
| 72°C | 7min | 1       |
| 20°C | 5min | 1       |

**NB: These PCR conditions have been optimized for high-throughput genotyping. Adaptation to small-scale may be required.**

## 2. Cre and Flp genotyping method

You will find the genotyping protocol in the publication:

[Highly-efficient, fluorescent, locus directed cre and FlpO deleter mice on a pure C57BL/6N genetic background.](#)

Birling MC, Dierich A, Jacquot S, Hérault Y, Pavlovic G.  
Genesis. 2012 Jun;50(6):482-9. doi: 10.1002/dvg.20826. Epub 2012 Mar 20.

